

SEQUENCE LISTING

<110> HILDEBRAND, DAVID
 RAO, SURYADEVARA S.

<120> SOYBEAN SELECTION SYSTEM BASED ON AEC-RESISTANCE

<130> 47100-222154

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<150> PCT/US04/020039

<151> 2004-06-23

<150> 60/483,103

<151> 2003-06-30

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<170> PatentIn Ver. 3.3

<210> 1

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<212> DNA

<213> Glycine max

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<223> a, c, g, t, unknown or other

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<221> MOD_RES

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<223> a, c, g, t, unknown or other

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Ala Ser Gly Thr Ser Ala Ile Val Ser Val Gly Thr Thr Gly Glu Ser
      35             40             45

Ala Thr Leu Asn His Asp Glu His Ala Asp Val Val Met Met Thr Leu
      50             55             60

Asp Leu Ala Asp Gly Arg Ile Pro Val Ile Ala Gly Thr Gly Ala Asn
      65             70             75             80

Ala Thr Ala Glu Ala Ile Ser Leu Thr Gln Arg Phe Asn Asp Ser Gly
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Ile Val Gly Cys Leu Thr Val Thr Pro Tyr Tyr Asn Arg Pro Ser Gln
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Glu Gly Leu Tyr Gln His Phe Lys Ala Ile Ala Glu His Thr Asp Leu
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Pro Gln Ile Leu Tyr Asn Val Pro Ser Arg Thr Gly Cys Asp Leu Leu
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Pro Glu Thr Val Gly Arg Leu Ala Lys Val Lys Asn Ile Ile Gly Ile
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 180 185 190
 Asp Phe Met Gln Leu Gly Gly His Gly Val Ile Ser Val Thr Thr Asn
 195 200 205
 Val Ala Ala Arg Asp Met Ala Gln Met Cys Lys Leu Ala Ala Glu Glu
 210 215 220
 His Phe Ala Glu Ala Arg Val Ile Asn Gln Arg Leu Met Pro Leu His
 225 230 235 240
 Asn Lys Leu Phe Val Glu Pro Asn Pro Ile Pro Val Lys Trp Ala Cys
 245 250 255
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 Ala Gly Leu Leu
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 35 40 45
 Phe Asp Leu Glu Ala Tyr Asp Asp Leu Val Asn Met Gln Ile Gly Gln
 50 55 60
 Gly Ala Glu Gly Val Ile Val Gly Gly Thr Thr Gly Glu Gly Gln Leu
 65 70 75 80
 Met Ser Trp Glu Glu His Ile Ile Leu Ile Ala His Thr Val Asn Cys
 85 90 95
 Phe Gly Gly Lys Ile Lys Val Ile Gly Asn Thr Gly Ser Asn Ser Thr
 100 105 110
 Arg Glu Ala Ile His Ala Thr Glu Gln Gly Phe Ala Val Gly Met His
 115 120 125

Ala Ala Leu His Ile Asn Pro Tyr Tyr Gly Lys Thr Ser Leu Asp Gly
 130 135 140

Met Val Ala His Phe Arg Ser Val Leu Ser Met Gly Pro Thr Ile Ile
 145 150 155 160

Tyr Asn Val Pro Ala Arg Thr Gly Gln Asp Ile Pro Pro His Val Ile
 165 170 175

Gln Thr Leu Ala Glu Ser Val Asn Leu Ala Gly Val Lys Glu Cys Val
 180 185 190

Gly Asn Asp Arg Ile Lys Gln Tyr Thr Asp Asp Gly Ile Val Val Trp
 195 200 205

Ser Gly Asn Asp Asp Gln Cys His Asp Ala Arg Trp Gly Tyr Gly Ala
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Thr Gly Val Val Ser Val Ala Ser Asn Leu Val Pro Gly Leu Met Arg
 225 230 235 240

Glu Leu Met Phe Gly Gly Val Asn Pro Thr Leu Asn Ser Lys Leu Leu
 245 250 255

Pro Leu Ile Asp Trp Leu Phe His Met Pro Asn Pro Ile Gly Leu Asn
 260 265 270

Thr Ala Leu Ala Gln Leu Gly Val Ile Arg Pro Val Phe Arg Leu Pro
 275 280 285

Phe Val Pro Leu Pro Val Asp Lys Arg Ile Glu Phe Ala Asn Leu Val
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Lys Glu Ile Gly Arg Glu His Phe Val Gly Asn Lys Val Val Glu Val
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<212> PRT

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 35 40 45

Lys Val Ile Gly Asn Thr Gly Ser Asn Ser Thr Arg Glu Ala Ile His
 50 55 60

Ala	Thr	Glu	Gln	Gly	Phe	Ala	Val	Gly	Met	His	Ala	Ala	Leu	His	Ile
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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
consensus sequence

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